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The following Listing of the Claims will replace all prior versions and all prior listings of the claims in the present application:

Listing of The Claims:

1-25. (Cancelled)

26. (Currently Amended) A method for inhibiting bacterial growth, comprising contacting bacteria <u>in vitro</u> with an amount of an inhibitor effective to reduce the activity of a polypeptide comprising the amino acid sequence of SEQ ID NO: 16, <u>wherein said inhibitor inhibits bacterial</u>

27-28. (Cancelled)

growth.

- 29. (Previously presented) The method of claim 26 wherein said inhibitor is selected from the group consisting of a small molecule, a peptidomimetic compound, and a bacterial growth inhibitory bacteriophage polypeptide.
- 30. (Previously Amended) The method of claim 26 wherein said inhibitor is a peptide synthesized by a recombinant expression system and purified, or artificially synthesized.

31-52. (Cancelled)

- 53. (Currently Amended) A method for inhibiting bacterial growth, comprising contacting a bacteria <u>in vitro</u> with an effective amount of an inhibitor <del>capable of decreasing</del> that decreases the activity of a polypeptide selected from the group consisting of:
  - a polypeptide comprising the amino acid sequence of SEQ ID NO: 2;

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- a polypeptide comprising the amino acid sequence of SEQ ID NO: 16; and

- a polypeptide comprising the amino acid sequence of SEQ ID NO: 18 wherein said inhibitor inhibits bacterial growth.

54-56. (Cancelled)

- 57. (Previously presented) The method of claim 53, wherein said inhibitor is selected from the group consisting of a small molecule, a peptidomimetic compound, and a bacterial growth inhibitory bacteriophage polypeptide.
- 58. (Previously presented) The method of claim 53, wherein said an inhibitor is a peptide synthesized by a recombinant expression system and purified, or artificially synthesized.
- 59. (Currently Amended) A method for inhibiting bacterial growth, comprising contacting a bacteria *in vitro* with an amount of an inhibitor effective to decrease the activity of a polypeptide selected from the group consisting of:
  - a DnaI polypeptide comprising at least 75% identity over 50 or more amino acids to the amino acid sequence of SEQ ID NO: 2;
  - a DnaI polypeptide comprising at least 85% similarity over 50 or more amino acids to the amino acid sequence of SEQ ID NO: 2;
  - a DnaI polypeptide comprising at least 75% identity over 50 or more amino acids to the amino acid sequence of SEQ ID NO: 16;
  - a DnaI polypeptide comprising at least 85% similarity over 50 or more amino acids to the amino acid sequence of SEQ ID NO: 16;
  - a DnaI polypeptide comprising at least 75% identity over 50 or more amino acids to the amino acid sequence of SEQ ID NO: 18;
  - a DnaI polypeptide comprising at least 85% similarity over 50 or more amino acids to the amino acid sequence of SEQ ID NO: 18; and

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fragments comprising an amino acid sequence having at least 50 contiguous amino acids from the amino acid of SEQ ID NO: 2; SEQ ID NO: 16; and SEQ ID NO: 18;

wherein said polypeptide has an activity selected from the group consisting of:

- a) directly interacting with bacteriophage 77 ORF 104 protein or a DnaI-binding fragment thereof in a manner that results in at least 10 fold reduction of <sup>3</sup> H-thymidine incorporation in a bacterial DNA replication assay relative to <sup>3</sup> H-thymidine incorporation in an assay lacking bacteriophage 77 ORF 104 protein or a DnaI-binding fragment thereof;
- b) directly interacting with bacteriophage 77 ORF 104 protein or a DnaI-binding fragment thereof in a manner that results in at least 10% inhibition of plasmid replication by bacteriophage 77 ORF 104 protein or a DnaI-binding fragment in a plasmid replication assay; and
- c) aiding in the loading of S. aureus DnaC helicase onto replicative primosomes wherein said inhibitor inhibits bacterial growth.

60-62 (Cancelled)

- 63. (Previously presented) The method of claim 59, wherein said inhibitor is selected from the group consisting of a small molecule, a peptidomimetic compound, and a bacterial growth inhibitory bacteriophage polypeptide.
- 64. (Previously presented) The method of claim 59, wherein said inhibitor is a peptide synthesized by a recombinant expression system and purified, or artificially synthesized.

65-66. (Cancelled)

67. (Currently amended) A method for inhibiting a bacterium bacterial growth, comprising contacting the bacterium bacteria *in vitro* with an inhibitor binding to an active domain of *S. aureus* DnaI, wherein said inhibitor inhibits bacterial growth.

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- 68. (Previously presented) The method of claim 67, wherein said active domain comprises amino acids selected from the group consisting of amino acids 1-313, amino acids 64-313, and amino acids 150-313 from SEQ ID NO: 2.
- 69. (Previously presented) The method of claim 67, wherein said inhibitor consists of an antibacterial agent inhibiting the biological activity of said *S. aureus* DnaI.
- 70. (Previously presented) The method of claim 69, wherein said biological activity is selected from the group consisting of:
  - a) directly interacting with bacteriophage 77 ORF 104 protein or a DnaI-binding fragment thereof in a manner that results in at least 10 fold reduction of <sup>3</sup> H-thymidine incorporation in a bacterial DNA replication assay relative to <sup>3</sup> H-thymidine incorporation in an assay lacking bacteriophage 77 ORF 104 protein or a DnaI-binding fragment thereof;
  - b) directly interacting with bacteriophage 77 ORF 104 protein or a DnaI-binding fragment thereof in a manner that results in at least 10% inhibition of plasmid replication by bacteriophage 77 ORF 104 protein or a DnaI-binding fragment in a plasmid replication assay; and
  - c) aids in the loading of S. aureus DnaC helicase onto replicative primosomes.
- 71. (Previously presented) The method of claim 67, wherein said binding inhibits *S. aureus* DnaI activity of aiding in the loading of *S. aureus* DnaC helicase onto replicative primosomes.
- 72. (Cancelled)
- 73. (Cancelled)
- 74. (Currently amended) A method for inhibiting bacterial DNA synthesis, comprising contacting a bacterium *in vitro* with an effective amount of an inhibitor <del>capable of decreasing</del> which decreases the activity of a polypeptide selected from the group consisting of:

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- a DnaI polypeptide comprising at least 75% identity over 50 or more amino acids to the amino acid sequence of SEQ ID NO: 2;

- a DnaI polypeptide comprising at least 85% similarity over 50 or more amino acids to the amino acid sequence of SEQ ID NO: 2;
- a DnaI polypeptide comprising at least 75% identity over 50 or more amino acids to the amino acid sequence of SEQ ID NO: 16;
- a DnaI polypeptide comprising at least 85% similarity over 50 or more amino acids to the amino acid sequence of SEQ ID NO: 16;
- a DnaI polypeptide comprising at least 75% identity over 50 or more amino acids to the amino acid sequence of SEQ ID NO: 18;
- a DnaI polypeptide comprising at least 85% similarity over 50 or more amino acids to the amino acid sequence of SEQ ID NO: 18; and
- fragments comprising an amino acid sequence having at least 50 contiguous amino acids from the amino acid of SEQ ID NO: 2; SEQ ID NO: 16; and SEQ ID NO: 18;

wherein said polypeptide has an activity selected from the group consisting of:

- a) directly interacting with bacteriophage 77 ORF 104 protein or a DnaI-binding fragment thereof in a manner that results in at least 10 fold reduction of <sup>3</sup> H-thymidine incorporation in a bacterial DNA replication assay relative to <sup>3</sup> H-thymidine incorporation in an assay lacking bacteriophage 77 ORF 104 protein or a DnaI-binding fragment thereof;
- b) directly interacting with bacteriophage 77 ORF 104 protein or a DnaI-binding fragment thereof in a manner that results in at least 10% inhibition of plasmid replication by bacteriophage 77 ORF 104 protein or a DnaI-binding fragment in a plasmid replication assay; and
- c) aiding in the loading of S. aureus DnaC helicase onto replicative primosomes,

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wherein said decrease in activity inhibitor inhibits bacterial DNA synthesis.